Research Article

Punctaporonins N–S, New Caryophyllene Sesquiterpenoids from *Cytospora* sp.

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Six new caryophyllene sesquiterpenoids, punctaporonins N–S (1–6), and three known ones, 6-hydroxypunctaporonins B (7), A (8), and E (9), have been isolated from solid cultures of *Cytospora* sp. The structures of 1–6 were elucidated primarily by NMR spectroscopy. The absolute configuration of 1 was assigned by X-ray crystallographic analysis of its S-MTPA ester. Compounds 2, 5, and 6 showed modest cytotoxicity against HeLa cells.

1. Introduction

Fungi are capable of producing a variety of bioactive secondary metabolites [1]. Since the secondary metabolism of fungi may be influenced by selection pressures exerted by other organisms and the environment in which they reside, those species thriving in unique and competitive niches are especially likely to produce bioactive natural products with diverse and interesting structural features [2, 3]. Based on this consideration and the documented success in finding new bioactive natural products from special types of fungi [4], we initiated chemical studies of the fungi inhabiting either the fruiting body and larvae of *Cordyceps sinensis* [5–10] or its surface soil [11–13]. As an extension, we also studied those species isolated from the soil samples that were collected on the Qinghai-Tibetan plateau at altitudes above 3,200 m, the environment in which *Cordyceps sinensis* was typically found. During the course of our continuing search for new bioactive natural products from this kind of fungal species, an ascomycetous fungus *Cytospora* sp. was isolated from a soil sample that was collected at Linzhi, Tibet, China. Our initial investigation of this fungus that was fermented in rice medium at 25°C led to the isolation of three antimicrobial caryophyllene-derived meroterpenoids [14, 15] and three

cytotoxic caryophyllene sesquiterpenoids [16]. In the current work, the fungus was refermented in the same solid culture medium at 15°C, and the HPLC chromatogram of the crude extract revealed the presence of additional components. Fractionation of an EtOAc extract afforded six new caryophyllene sesquiterpenoids, which we named punctaporonins N–S (1–6), and three known ones, 6-hydroxypunctaporonins B (7), A (8), and E (9) (Figure 1) [17]. Details of the isolation, structure elucidation, and cytotoxicity of these compounds are described herein.

2. Materials and Methods

2.1. General Experimental Procedures. Optical rotations were measured on a Perkin-Elmer 241 polarimeter, and UV data were recorded on a Shimadzu Biospec-1601 spectrophotometer. IR data were recorded using a Nicolet Magna-IR 750 spectrophotometer. ¹H and ¹³C NMR data were acquired with Varian Mercury-400, Varian Mercury-500, and Varian Mercury-600 MHz spectrometers using solvent signals (acetone-*d*₆: δ_H 2.05/δ_C 29.8, 206.1; pyridine-*d*₅: δ_H 7.21, 7.58, 8.73) as references. The HMQC and HMBC experiments were optimized for 145.0 and 8.0 Hz, respectively. ESIMS data and

FIGURE 1: Structures of compounds 1–9.

HRESIMS data were obtained using an Agilent Accurate-Mass-Q-TOF LC/MS 6520 instrument equipped with an electrospray ionization (ESI) source. The fragmentor and capillary voltages were kept at 125 and 3500 V, respectively. Nitrogen was supplied as the nebulizing and drying gas (300°C). The flow rate of the drying gas and the pressure of the nebulizer were 10 L/min and 10 psi, respectively. All MS experiments were performed in positive ion mode. Full-scan spectra were acquired over a scan range of m/z 100–1000 at 1.03 spectra/s. All solvents used were of analytical grade. Column chromatography was performed with silica gel (100–200 or 200–300 mesh, Qingdao Marine Chemical Inc., China) and Sephadex LH-20 (GE, USA). Semipreparative HPLC was performed on an Agilent 1260 G1311C isopump equipped with a G1365D MWD detector and an Agilent Zorbax SB-C₁₈ column (5 μ m; 9.40 mm $\times$ 250 mm).

2.2. Fungal Material and Fermentation. The isolation, identification, and fermentation of the fungus *Cytospora* sp. were the same as those we previously described [14–16], except the incubation temperature was changed to 15°C.

2.3. Extraction and Isolation. The fermented material was extracted with EtOAc (4 $\times$ 1.0 L), and the organic solvent was evaporated to dryness under vacuum to afford a crude extract (4.2 g). The extract was fractionated by silica gel VLC using petroleum ether–EtOAc gradient elution. The fraction (100 mg) eluted with 35% EtOAc was separated by Sephadex LH-20 column chromatography (CC) eluting with 1:1 CH₂Cl₂–MeOH. The resulting subfractions were combined and further purified by semipreparative RP HPLC (59% MeOH in H₂O for 35 min; 2 mL/min) to afford punctaporonins N (1; 4.5 mg, t_R 21.22 min), O (2; 3.5 mg, t_R 31.33 min), P (3; 2.0 mg, t_R 14.05 min), and R (5; 0.8 mg, t_R 24.22 min). The fraction (50 mg) eluted from the silica gel column with 45% EtOAc was purified by RP HPLC (35% CH₃CN in H₂O for 20 min; 2 mL/min) to afford punctaporonin Q (4; 5.0 mg; t_R 16.44 min). The fraction (35 mg) eluted from the silica gel column with 40% EtOAc was purified by RP HPLC (40% CH₃CN in H₂O for 20 min; 2 mL/min) to afford punctaporonin S (6; 2.0 mg; t_R

17.02 min). The fraction (150 mg) eluted from the silica gel column with 50% EtOAc was fractionated again by Sephadex LH-20 CC eluting with 1:1 CH₂Cl₂–MeOH. Purification of the resulting subfractions with different gradients afforded 6-hydroxypunctaporonins B (7; 10.0 mg, t_R 23.61 min; 50% MeOH in H₂O for 30 min), A (8; 8.4 mg, t_R 16.12 min; same gradient as in purification of 7), and E (9; 3.5 mg, t_R 14.83 min; same gradient as in purification of 7).

2.4. Identification

Punctaporonin N (1). White powder (MeOH); $[\alpha]_D^{25}$ –73 (c 0.1, MeOH); UV (MeOH) λ_{max} (log ϵ) 198 (3.40) nm; IR (neat) ν_{max} 3317 (br), 2936, 2870, 1736, 1714, 1463, 1372, 1241, 1095, 1037 cm^{–1}; ¹H, ¹³C NMR, HMBC, and NOESY data see Table 1; (+)–HR–ESI–MS m/z 305.1719 [M + Na]⁺ (calcd. for C₁₆H₂₆O₄Na, 305.1723).

Punctaporonin O (2). Colorless oil (MeOH); $[\alpha]_D^{25}$ –47 (c 0.1, MeOH); UV (MeOH) λ_{max} (log ϵ) 202 (3.58) nm; IR (neat) ν_{max} 3263 (br), 2932, 2875, 1713, 1463, 1361, 1181, 1089, 1058, 1029 cm^{–1}; ¹H and ¹³C NMR data see Table 2; (+)–HR–ESI–MS m/z 305.1721 [M + Na]⁺ (calcd. for C₁₆H₂₆O₄Na, 305.1723).

Punctaporonin P (3). Colorless oil (MeOH); $[\alpha]_D^{25}$ –71 (c 0.1, MeOH); UV (MeOH) λ_{max} (log ϵ) 203 (3.55) nm; IR (neat) ν_{max} 3262 (br), 2958, 2935, 2872, 1735, 1717, 1372, 1240, 1089, 1021 cm^{–1}; ¹H and ¹³C NMR data see Table 2; (+)–HR–ESI–MS m/z 333.1669 [M + Na]⁺ (calcd. for C₁₇H₂₆O₅Na, 333.1672).

Punctaporonin Q (4). Colorless oil (MeOH); $[\alpha]_D^{25}$ –37 (c 0.1, MeOH); UV (MeOH) λ_{max} (log ϵ) 203 (3.60) nm; IR (neat) ν_{max} 3438 (br), 2957, 2936, 1730, 1715, 1381, 1249, 1031 cm^{–1}; ¹H and ¹³C NMR data see Table 2; (+)–HR–ESI–MS m/z 333.1676 [M + Na]⁺ (calcd. for C₁₇H₂₆O₅Na, 333.1672).

Punctaporonin R (5). Colorless oil (MeOH); $[\alpha]_D^{25}$ –30 (c 0.1, MeOH); UV (MeOH) λ_{max} (log ϵ) 200 (3.22) nm; IR (neat) ν_{max} 3256 (br), 2957, 2934, 2869, 1711, 1462, 1365, 1102, 1057, 1031, 984 cm^{–1}; ¹H and ¹³C NMR data see

TABLE 1: NMR data (400 MHz, Acetone- d_6) for punctaporonin N (**1**).

Position	δ_C , mult.	δ_H (J in Hz)	HMBC ^a	NOESY
1	140.0, qC			
2	39.8, CH	3.39, dd (11.9, 8.2)	1, 3, 5, 11, 12	13
3a	34.0, CH ₂	2.15, dd (11.9, 9.8)	1, 2, 4, 13, 14	
3b		1.49, dd (9.8, 8.2)	2, 4, 5, 13	13
4	41.0, qC			
5	82.7, qC			
6	71.4, CH	3.86, br d (4.7)		13, 16
7a	42.5, CH ₂	2.63, dd (16.3, 4.7)	5, 6, 15	
7b		1.64, dt (16.3, 2.0)	6, 8, 9, 15	
8	78.4, qC			
9	141.9, CH	5.70, dt (13.0, 2.0)	7, 10, 11	
10	124.5, CH	5.62, d (13.0)	1, 9	
11	125.9, CH	5.88, s	1, 2, 9, 12	12b
12a	64.8, CH ₂	4.19, d (12.0)	1, 11	
12b		3.93, d (12.0)	1, 2, 11	11, 13
13	24.6, CH ₃	1.23, s	3, 4, 5, 14	2, 3b, 6, 12b
14	24.1, CH ₃	1.07, s	3, 4, 5, 13	OH-5
15	26.0, CH ₃	1.10, s	7, 8, 9	16
16	48.7, CH ₃	3.26, s	8	6, 15
OH-5		4.53, br s		14
OH-6		4.67, br s		
OH-12		3.14, br s		

^aHMBC correlations, optimized for 8 Hz, are from proton(s) stated to the indicated carbon.

Table 3; (+)-HR-ESI-MS m/z 305.1717 [$M + Na$]⁺ (calcd. for C₁₆H₂₆O₄Na, 305.1723).

Punctaporonin S (6). Colorless oil (MeOH); $[\alpha]_D^{25}$ -84 (c 0.1, MeOH); UV (MeOH) λ_{max} (log ϵ) 202 (3.56) nm; IR (neat) ν_{max} 3486 (br), 2953, 2938, 2871, 1736, 1374, 1252, 1040, 990 cm⁻¹; ¹H and ¹³C NMR data see Table 3; (+)-HR-ESI-MS m/z 375.1781 [$M + Na$]⁺ (calcd. for C₁₉H₂₈O₆Na, 375.1778).

6-Hydroxypunctaporonin B (7). ¹H, ¹³C NMR, and the MS data were consistent with literature values [17].

6-Hydroxypunctaporonin A (8). ¹H, ¹³C NMR, and the MS data were consistent with literature values [17].

6-Hydroxypunctaporonin E (9). ¹H, ¹³C NMR, and the MS data were consistent with literature values [17].

2.5. Preparation of (S)-MTPA Ester (1a). A sample of **1** (2.0 mg, 0.007 mmol) was dissolved in pyridine (2.0 mL) in a 10 mL round-bottomed flask. (R)-MTPA Cl (2.0 μ L, 0.011 mmol) were quickly added, and the flask was sealed and all contents were stirred at room temperature for 12 h. The mixture was evaporated to dryness and purified by semipreparative HPLC (90% CH₃OH in H₂O for 20 min; 2 mL/min) to afford **1a** (2.2 mg, t_R 15.72 min): colorless platelets; ¹H NMR (pyridine- d_5 , 500 MHz) δ 6.16 (1H, s, H-11), 6.00 (1H, d, J = 13 Hz, H-9), 5.75 (1H, d, J = 13 Hz, H-10),

5.55 (1H, d, J = 12 Hz, H-12a), 5.46 (1H, d, J = 12 Hz, H-12b), 4.37 (1H, s, H-6), 3.35 (3H, s, H₃-16), 3.09 (1H, d, J = 10 Hz, H-2), 2.34 (1H, t, J = 10 Hz, H-3a), 2.11 (1H, d, J = 16 Hz, H-7a), 1.54 (1H, t, J = 10 Hz, H-3b), 1.35 (1H, d, J = 16 Hz, H-7b), 1.37 (3H, s, H₃-13), 1.37 (3H, s, H₃-14), 1.18 (3H, s, H₃-15).

2.6. X-Ray Crystallographic Analysis of 1a [18]. Upon crystallization from MeOH/H₂O (10:1) using the vapor diffusion method, colorless crystals were obtained for **1a**, and a crystal (0.90 $\times$ 0.27 $\times$ 0.08 mm³) was separated from the sample and mounted on a glass fiber, and data were collected using a Rigaku Saturn CCD area detector with graphite-monochromated Mo K α radiation, λ = 0.71073 Å at 173(2) K. Crystal data: C₂₆H₃₃F₃O₆·H₂O, M = 516.54, space group orthorhombic, $P2_12_12_1$; unit cell dimensions a = 6.9639 (14) Å, b = 17.211 (3) Å, c = 21.948 (5) Å, V = 2630.5 (9) Å³, Z = 4, D_{calcd} = 1.304 mg/m³, μ = 0.107 mm⁻¹, $F(000)$ = 1096. The structure was solved by direct methods using SHELXL-2016 [19] and refined using full-matrix least-squares difference Fourier techniques. All nonhydrogen atoms were refined with anisotropic displacement parameters, and all hydrogen atoms were placed in idealized positions and refined as riding atoms with the relative isotropic parameters. Absorption corrections were applied with the Siemens Area Detector Absorption Program (SADABS) [20]. The 14274 measurements yielded 4799 independent reflections after equivalent data were averaged, and Lorentz and polarization corrections were applied. The final refinement gave R_1 = 0.0811 and wR_2 = 0.1232 [$I > 2\sigma(I)$].

TABLE 2: NMR spectroscopic data for punctaporonins O–Q (2–4) in acetone- d_6 .

Position	2		3		4	
	δ_C^a , mult.	δ_H^b (J in Hz)	δ_C^a , mult.	δ_H^b (J in Hz)	δ_C^a , mult.	δ_H^b (J in Hz)
1	55.3, qC		55.6, qC		53.6, qC	
2	47.1, CH	2.54, dd (13.0, 8.0)	47.2, CH	2.32, dd (12.8, 7.8)	45.6, CH	2.35, dd (12.5, 7.9)
3a	35.5, CH ₂	2.02, dd (13.0, 8.0)	35.5, CH ₂	2.08, dd (12.8, 8.9)	36.3, CH ₂	2.13, dd (12.5, 9.2)
3b		1.46, t (8.0)		1.52, dd (8.9, 7.8)		1.46, dd (9.2, 7.9)
4	42.8, qC		42.9, qC		42.7, qC	
5	80.6, qC		80.4, qC		81.7, qC	
6	68.9, CH	4.09, dd (10.0, 5.6)	68.7, CH	4.06, ddd (10.3, 7.9, 5.6)	68.9, CH	4.17, dd (10.0, 5.9)
7a	41.5, CH ₂	1.86, dd (13.5, 5.6)	41.2, CH ₂	1.77, dd (14.0, 5.6)	40.3, CH ₂	1.90, dd (13.5, 5.9)
7b		1.56, dd (13.5, 10.0)		1.62, dd (14.0, 10.3)		1.64, dd (13.5, 10.0)
8	51.0, qC		50.0, qC		50.9, qC	
9	97.3, CH	3.60, d (2.0)	89.4, CH	5.11, d (2.5)	86.5, CH	4.07, d (2.5)
10	127.9, CH	5.90, dd (6.0, 2.0)	126.6, CH	5.67, dd (5.9, 2.5)	130.6, CH	5.68, dd (6.0, 2.5)
11	143.4, CH	5.74, d (6.0)	145.3, CH	5.88, d (5.9)	141.2, CH	5.92, d (6.0)
12a	62.2, CH ₂	3.68, d (11.4)	62.0, CH ₂	3.73, d (11.2)	65.6, CH ₂	4.57, d (11.6)
12b		3.40, d (11.4)		3.48, dd (11.2, 8.0)		4.37, d (11.6)
13	23.4, CH ₃	1.11, s	23.4, CH ₃	1.19, s	23.3, CH ₃	1.10, s
14	24.1, CH ₃	1.08, s	24.0, CH ₃	1.12, s	23.7, CH ₃	1.05, s
15	24.6, CH ₃	1.05, s	24.3, CH ₃	1.10, s	26.2, CH ₃	0.84, s
16	57.5, CH ₃	3.33, s				
OH-5		4.82, br s		4.88, s		
OH-6		5.39, br s		5.37, d (7.9)		
OH-9						3.29, s
OH-12		3.25, br s		3.44, d (8.0)		
Ac-9			21.3, CH ₃ 170.4, qC	2.03, s		
Ac-12					21.0, CH ₃ 171.0, qC	1.93, s

^aRecorded at 100 MHz. ^bRecorded at 500 MHz.

2.7. MTS Assay. The MTS assay method was the same as that we previously described [16, 21, 22].

3. Results and Discussion

Punctaporonin N (**1**) was obtained as a white powder with a molecular formula of C₁₆H₂₆O₄ (four degrees of unsaturation), established by HRESIMS. Analysis of its NMR spectroscopic data (Table 1) revealed structural similarity to the coisolated known compound 6-hydroxypunctaporonin B (**7**) [17], except for the presence of an *O*-methyl group (δ_H/δ_C 3.26/48.7) attached to C-8, rather than a free hydroxy group as found in **7**. This observation was supported by an HMBC correlation from the *O*-methyl proton signal (H₃-16) to the oxygenated sp³ quaternary carbon at 78.4 ppm (C-8). Therefore, **1** was proposed as the C-8 methyl ether of **7**.

The relative configuration of **1** was determined by analysis of the ¹H–¹H coupling constants and NOESY data (Figure 2). The coupling constant of 13.0 Hz between H-9 and H-10, which was exactly the same as in **7**, suggested the *Z*-geometry for the C-9/C-10 olefin [17]. The C-1/C-11 olefin was deduced to be *E*-configuration based on NOESY correlation of H-11

with H-12b. NOESY correlations of H₃-13 with H-2, H-3b, and H-6 and H-6 with H₃-16 indicated that these protons are all on the same face of the ring system, whereas that of H₃-14 with OH-5 was used to place them on the opposite face, thereby establishing the relative configuration of **1**.

The modified Mosher method was tried to apply assigning the C-6 absolute configuration of **1**. However, treatment of **1** with (*R*) or (*S*)-MTPA Cl to form the C-6 MTPA ester was unsuccessful; instead the reaction products C-12 *S*-MTPA ester (**1a**) and C-12 *R*-MTPA ester (**1b**) were generated. In order to establish the absolute configuration of **1**, one of the reaction products C-12 *S*-MTPA ester (**1a**) was purified by HPLC to obtain a single crystal, and its perspective ORTEP plot is shown in Figure 3. Although the X-ray diffraction was collected by Mo K α radiation, which only gave the relative configuration for **1a**, the absolute configuration of the chiral center at C-18 generated from (*R*)-MTPA Cl had been assigned to be *S*. Therefore, the absolute configuration of **1** was established as 2*R*, 5*S*, 6*S*, and 8*S* by single-crystal X-ray crystallographic analysis of its C-12 *S*-MTPA ester (**1a**).

Punctaporonin O (**2**) was assigned the same molecular formula C₁₆H₂₆O₄ as **1** by HRESIMS. Its NMR data (Table 2)

TABLE 3: NMR spectroscopic data for punctaporonins R (5) and S (6) in acetone- d_6 .

Position	5	6
	δ_C^a , mult.	δ_C^c , mult.
	δ_H^b (J in Hz)	δ_H^b (J in Hz)
1	51.2, qC	53.6, qC
2	43.8, CH	39.2, CH
3a	35.8, CH ₂	37.8, CH ₂
3b		
4	42.1, qC	42.2, qC
5	80.2, qC	82.6, qC
6	69.3, CH	69.0, CH
7a	42.9, CH ₂	42.4, CH ₂
7b		
8	53.6, qC	48.8, qC
9	149.2, CH	150.0, CH
10	126.8, CH	126.8, CH
11	90.9, CH	81.3, CH
12a	57.3, CH ₂	63.6, CH ₂
12b		
13	23.3, CH ₃	22.9, CH ₃
14	24.2, CH ₃	23.6, CH ₃
15	27.3, CH ₃	27.9, CH ₃
16	60.7, CH ₃	
OH-5	5.29, br s	
OH-6	5.18, d (7.3)	3.69, (5.5)
OH-12	3.09, d (7.0)	
Ac-11		20.8, CH ₃
		170.1, qC
Ac-12		21.0, CH ₃
		170.9, qC

^aRecorded at 150 MHz. ^bRecorded at 500 MHz. ^cRecorded at 100 MHz.

FIGURE 2: Key NOESY correlations for punctaporonin N (1).

were nearly identical to those of the coisolated known compound 6-hydroxypunctaporonin A (8) [17], except that the exchangeable proton at 4.18 ppm (OH-9) was replaced by a methyl group (δ_H/δ_C 3.33/57.5). An HMBC correlation from the O-methyl proton signal (H₃-16) to the oxymethine carbon at 97.3 ppm (C-9) suggested that 2 is the C-9 methyl ether of 8. The absolute configuration of 2 was proposed

as shown by analogy to 8, which was also supported by the nearly identical specific rotation values recorded for both compounds [17].

The molecular formula of punctaporonin P (3) was determined to be C₁₇H₂₆O₅ (five degrees of unsaturation) by HRESIMS, which is 28 mass units more than that of 2, corresponding to an extra carbonyl group. The NMR data of 3 (Table 2) were similar to those of 2, except that the oxymethine proton at 3.60 ppm (H-9) was significantly downfield to 5.11 ppm in 3. In addition, the O-methyl group attached to C-9 was replaced by an acetyl unit (δ_H/δ_C 2.03/21.3, 170.4), indicating that the C-9 oxygen of 3 was acetylated. An HMBC correlation from the downfield oxymethine proton (H-9) to the carboxylic carbon at 170.4 ppm indicated that 3 is the C-9 monoacetate of 8, with its configuration similarly deduced by analogy to compounds 1 and 2.

Punctaporonin Q (4) gave a pseudomolecular ion [M + Na]⁺ peak by HRESIMS, consistent with the molecular formula C₁₇H₂₆O₅ (five degrees of unsaturation), which is the same as 3. Its NMR spectra showed resonances similar to those of 3, indicating that 4 is also a monoacetate of 8, but with a different position for acetylation. Specifically, the ¹H NMR chemical shifts of H₂-12 in 4 (δ_H 4.37 and 4.57,

FIGURE 3: Thermal ellipsoid representation of **1a**.

Table 2) were significantly downfield compared to those in **3** (δ_{H} 3.48 and 3.73), suggesting that OH-12 was acetylated. The observation was supported by an HMBC correlation from H₂-12 to the carboxylic carbon of the acetyl unit (δ_{C} 171.0).

The elemental composition of punctaporonin R (**5**) was established as C₁₆H₂₆O₄ (four degrees of unsaturation) by HRESIMS, which has one more CH₂ unit than another coisolated known compound, 6-hydroxypunctaporonin E (**9**) [17]. Comparison of the NMR data of **5** (Table 3) and **9** readily identified **5** as the C-11 methyl ether ($\delta_{\text{H}}/\delta_{\text{C}}$ 3.19/60.7) of the latter and having the same configuration as its precedent.

Punctaporonin S (**6**) was assigned the molecular formula C₁₉H₂₈O₆ (six degrees of unsaturation) by HRESIMS. Its NMR spectra showed structural characteristics similar to those of **5**, but the ¹H NMR chemical shifts of H-11 (δ_{H} 5.70) and H₂-12 (δ_{H} 4.06 and 4.88) were significantly downfield compared to **5** (H-11: δ_{H} 3.62; H₂-12: δ_{H} 3.65 and 4.06), indicating that both OH-11 and OH-12 are acetylated, which were supported by relevant HMBC correlations. Therefore, **6** was assigned as the diacetate of the known compound **9**, with its configuration deduced as shown.

To verify that the new metabolites **1–6** are authentic natural products, a portion of the freeze-dried fermented rice substrate was extracted with distilled, HPLC grade acetone, and the resulting extract was subjected to RP HPLC analysis using distilled, HPLC grade H₂O and MeOH as solvents. Compounds **1–6** were identified on the HPLC chromatogram of the crude extract by comparison of their retention times with the pure compounds, indicating that **1–6** are indeed naturally occurring metabolites.

Compounds **2**, **5**, and **6** showed modest cytotoxicity against HeLa (cervical epithelium) cells, showing IC₅₀ values of 16.6, 10.4, and 47.4 μM , respectively, while the positive control

cisplatin showed an IC₅₀ value of 7.6 μM . The remaining compounds did not show noticeable cytotoxic effects against HeLa cells (IC₅₀ > 100 μM).

4. Conclusion

In summary, six new caryophyllene sesquiterpenoids, punctaporonins N–S (**1–6**), and three known ones, 6-hydroxypunctaporonins B (**7**), A (**8**), and E (**9**), have been isolated from solid cultures of *Cytospora* sp. The structures of **1–6** were elucidated primarily by NMR spectroscopy. The absolute configuration of **1** was assigned by X-ray crystallographic analysis of its S-MTPA ester. Compounds **2**, **5**, and **6** showed modest cytotoxicity against HeLa cells. These results implied that the fungi isolated from those unique and competitive niches could prove to be valuable sources of new bioactive natural products.

Conflicts of Interest

The authors have declared that there are no conflicts of interest.

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